WHAT IS CLAIMED IS:

1	1. A method of screening an individual for increased risk of low		
2	folate status, said method comprising detecting a mutation in a human glutamate		
3	carboxypepidase II (GCPII) gene in a biological sample from said individual, wherein		
4	detection of the mutation is indicative of decreased ability to hydrolyse a terminal		
5	glutamate residue of a folypoly-γ-glutamate, which decreased ability is associated with		
6	low folate status.		
1	2. The method of claim 1, wherein the mutation is a single nucleotide		
2	polymorphism.		
4	porymorpinsm.		
1	3. The method of claim 3, wherein the single nucleotide		
2	polymorphism causes an amino acid substitution of H475Y.		
1	4. A method of claim 1 wherein the mutation is detected by		
2	(a) amplifying the GCPII gene, or a portion thereof containing the		
3	mutation, with a set of primers to provide an amplified product,		
4	(b) sequencing the amplified product to obtain a sequence, and		
5	(c) comparing the sequence of the amplified product with a known		
6	sequence of a wild-type GCPII gene,		
7	wherein a difference between the sequence of the amplified product and		
8	the sequence of the wild-type GCPII gene indicates the presence of a mutation.		
1	5. A method of claim 4, wherein said amplification is by polymerase		
2	chain reaction.		
1	6. A method of claim 4, wherein said sequencing is performed by		
2	detecting the incorporation of a nucleotide into a strand complementary to a template		
3	strand by detecting the presence of a pyrophosphate released from the incorporated		
4	nucleotide.		
1	7. A method of claim 1 wherein the mutation is detected by		
2	(a) amplifying exon 13 of the GCPII gene with a set of primers to		
3	provide an amplified product,		
4	(b) sequencing the amplified product to obtain a sequence, and		

5		(c) comparing the sequence of the amplified product with a known	
6	sequence of exon 13 of a wild-type GCPII gene,		
7	wherein a difference between the sequence of the amplified product and		
8	the sequence of the v	wild-type GCPII gene indicates the presence of a mutation.	
1	8.	A method of claim 7, wherein said primers are	
2	5'-CATTCTGGTAG	GGAATT TAGCA-3' and 5'-AAACACCACCTATGTTTAACA-3'.	
1	9.	A method of claim 7, wherein said amplification is by polymerase	
2	chain reaction.		
1	10.	A method of claim 7, wherein said sequencing is performed by	
2	detecting the incorporation of a nucleotide into a strand complementary to a template		
3	strand by detecting the presence of a pyrophosphate released from the incorporated		
4	nucleotide.		
1	11.	A method of claim 1, wherein said mutation is detected by	
2	hybridizing DNA fro	om said individual to a test nucleic acid under stringent conditions.	
1	12.	A method of claim 11, wherein either said DNA from said	
2	individual or said test nucleic acid is immobilized on a solid support.		
1	13.	A method of claim 1, wherein said mutation is detected by	
2		(a) amplifying exon 13 said GCPII gene,	
3		(b) subjecting said amplified exon 13 to digestion by restriction	
4	enzymes,		
5		(c) separating the resulting restriction products to form a pattern of	
6	restriction fragment lengths, and		
7		(d) comparing the pattern of restriction fragment lengths to a	
8	pattern of restriction fragment lengths formed by subjecting amplified exon 13 of a wild-		
9	type GCPII gene to the same restriction enzymes.		
1	14.	A method of claim 13, wherein said separation of the restriction	
2	products is by gel el	ectrophoresis.	

15.

A method of claim 13, wherein the restriction enzyme is AccI.

1			
1	16. A method of claim 15, wherein the pattern of restriction fragments		
2	of exon 13 of the GCPII gene of the individual shows restriction fragments selected from		
3	the group consisting of: 141 bases and 103 bases.		
1	17. A method of claim 1, wherein said mutation is detected by		
2	specifically binding an antibody to a truncated product of the GCPII gene, wherein the		
3	specific binding of the antibody to the truncated gene product is indicative of a mutation		
4	impairing the ability of the GCPII gene product to digest a dietary folate.		
1	18. A method of claim 17, wherein detection of said specific binding of		
2	said antibody and said truncated gene product is by ELISA.		
1	A method of screening an individual for increased risk of low		
2	folate status comprising		
3	(a) performing reverse transcriptase-PCR on mRNA from intestinal cells		
4	of the individual to amplify products of a GCPII gene, and		
5	(b) determining the ratio of a variant product in which 93 bases of exon 18		
6	are deleted to a normal product of the GCPII gene,		
7	wherein a ratio of the variant form to the normal form greater than 1:3		
8	indicates the individual is at increased risk of low folate status.		
1	20. A mutation in a GCPII gene which impairs the ability of a product		
2	of the gene to hydrolyse a conjugated folate to release folic acid compared to a product of		
3	a wild-type GCPII gene.		
1	21. A mutation of claim 20, wherein the ability of a product of the gen		
2	to hydrolyse a conjugated folate is reduced by 20 percent or more compared to a product		
3	of a wild-type GCPII gene.		
1	22. A mutation of claim 20, wherein the mutation is a 93-base deletion		
2	resulting from the elimination of exon 18.		
1	23. The mutation of claim 20, wherein the mutation is a single		
2	nucleotide polymorphism.		

polymorphism causes an amino acid substitution of: H475Y.

The mutation of claim 23, wherein the single nucleotide

24.

30.

1

2

A kit of claim 28, further comprising an AccI restriction enzyme.